

GenCore version 4.5
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OM nucleic - nucleic search, using sw model

Run on: March 16, 2000, 15:37:36 ; Search time 151.92 Seconds
(without alignments)
5342.438 Million cell updates/sec

Title: US-09-211-755-1
Perfect score: 3244
Sequence: 1 TGACCTCGGGCAGGTCCTG.....CTTGCAAAAAAAAAAAAA 3244

Scoring table: IDENTITY_NUC

Searched: 311585 seqs, 125096042 residues

Database : N_Geneseq_36:*

Word size : 0

Number of hits that pass the threshold : 623170

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

| Result No. | Score | Query Match | Length | DB ID | Description |
|------------|-------|-------------|--------|----------|--------------------|
| 1 | 317.2 | 9.8 | 2924 | 1 V10267 | Human GABA-BR1b re |
| 2 | 315.2 | 9.7 | 2620 | 1 V10265 | Human GABA-BR1a/b |
| 3 | 312.8 | 9.6 | 2837 | 1 V10266 | Rat GABA-BR1b rece |
| 4 | 306 | 9.4 | 4376 | 1 V10264 | Rat GABA-BR1a rece |
| 5 | 286.8 | 8.8 | 314 | 1 V88919 | EST clone HW456. N |
| 6 | 177 | 5.5 | 361 | 1 X51891 | Human secreted pro |
| 7 | 45.4 | 1.4 | 2176 | 1 Q11726 | Human secreted pro |
| 8 | 45.4 | 1.4 | 397 | 1 X51965 | Human secreted pro |
| 9 | 43.6 | 1.3 | 15872 | 1 Q10613 | Rianodin receptor |
| 10 | 42.4 | 1.3 | 273 | 1 V89458 | EST clone CO337. N |
| 11 | 42.4 | 1.3 | 114955 | 1 X53491 | Human adenosine A1 |
| 12 | 41.8 | 1.3 | 1312 | 1 Q73383 | Human CCAAT/Enhanc |
| 13 | 41.8 | 1.3 | 1312 | 1 T44325 | DNA encoding CCAAT |
| 14 | 41.6 | 1.3 | 8438 | 1 Q73500 | DNA encoding Pseud |
| 15 | 41.6 | 1.3 | 2823 | 1 T35233 | Natural killer lyt |
| 16 | 41.6 | 1.3 | 2888 | 1 T53268 | Streptomyces prist |
| 17 | 41 | 1.3 | 2618 | 1 T16710 | Metabotropic gluta |
| 18 | 41 | 1.3 | 2619 | 1 T29408 | Human metabotropic |
| 19 | 41 | 1.3 | 4131 | 1 T89290 | Dogfish shark kidn |
| 20 | 40.6 | 1.3 | 3865 | 1 Q37101 | Bovine transglutam |
| 21 | 40.6 | 1.3 | 10596 | 1 Q51731 | Plasmod pcisEBON f |
| 22 | 40.6 | 1.3 | 10596 | 1 T40348 | Plasmod pcisEBON f |
| 23 | 40.6 | 1.3 | 9600 | 1 V21683 | Vector plasmid pCM |
| 24 | 40.6 | 1.3 | 10596 | 1 X15650 | Nucleotide sequenc |
| 25 | 40.4 | 1.2 | 1080 | 1 T68789 | Melanocortin-3 rec |
| 26 | 40.4 | 1.2 | 1080 | 1 V06399 | Human melanocortin |
| 27 | 40.4 | 1.2 | 1080 | 1 V62351 | Melanocortin-3 rec |
| 28 | 40.4 | 1.2 | 1080 | 1 Q19363 | Human MC3 DNA. Pro |
| 29 | 40.4 | 1.2 | 1648 | 1 Q25532 | Sequence of genom |
| 30 | 40.2 | 1.2 | 1158 | 1 V06396 | Maize optimised ge |
| 31 | 40.2 | 1.2 | 799 | 1 V55831 | Nucleotide sequenc |
| 32 | 39.8 | 1.2 | 15377 | 1 Q25975 | MH mutant porcine |
| 33 | 39.4 | 1.2 | 2006 | 1 Q50147 | Phospholipase D-p |
| 34 | 39.2 | 1.2 | 5006 | 1 T61381 | Parathyroid calcu |
| 35 | 39.2 | 1.2 | 3809 | 1 T61382 | Parathyroid calcu |
| 36 | 39.2 | 1.2 | 1041 | 1 T77781 | Nuclear mitotic ap |
| 37 | 39.2 | 1.2 | 1092 | 1 T78310 | E6AP-binding prote |
| 38 | 39.2 | 1.2 | 4000 | 1 T86166 | Nucleotide sequenc |
| 39 | 39.2 | 1.2 | 5006 | 1 T95858 | Human parathyroid |

40 39.2 1.2 3809 1 T95859 Human parathyroid
41 39.2 1.2 5006 1 V26963 Human parathyroid
42 39.2 1.2 3809 1 V26964 Human parathyroid
43 39.2 1.2 5006 1 V82484 Human parathyroid
44 39.2 1.2 3809 1 V82485 Human parathyroid
45 39 1.2 1312 1 T93367 Mouse thrombin rec

ALIGNMENTS

RESULT 1
V10267
ID V10267 standard; cDNA to mRNA; 2924 BP.
AC V10267;
DT 03-JUN-1998 (first entry)
DE Human GABA-BR1b receptor: cDNA.
KW Gamma-aminobutyric acid; GABA-BR1a/b receptor; human; brain; agonist;
KW inhibitory neurotransmitter; peripheral nervous system; antagonist;
KW treatment; dementia; depression; anxiety; bronchial inflammation; asthma;
KW epilepsy; cognitive function; ds.
OS Homo sapiens.
FH Key Location/Qualifiers
FT 169..2703
FT /*tag= a
FT /product= GABA-BR1b
PN WO9746675-A1.
PD 11-DEC-1997.
PF 19-MAR-1997; E01370.
PR 22-NOV-1996; US-756091.
PR 30-MAY-1996; US-655716.
PA (NOVS) NOVARTIS AG.
PI Bettler B, Bittiger H, Froestl W, Kaupmann K, Mickel SJ;
DR WPI: 98-042183/04.
DR P-FSDB: W40119.
PT Purified GABA-B receptor or receptor protein - and antagonists of
these which may be useful in treating nervous system disorders
PS Claim 3; Page 79-86; 108pp; English.
CC This cDNA sequence encodes a novel human GABA-B receptor protein,
CC GABA-BR1b. GABA (gamma-aminobutyric acid) is the major inhibitory
CC neurotransmitter found in the brain and peripheral nervous system
CC and this receptor may be used for the identification of GABA-B
CC receptor agonists and antagonists. Such proteins may be used in
CC treatment of dementia, depression, anxiety, epilepsy, spasticity,
CC bronchial inflammation or asthma or to improve cognitive function.
CC GABA-B receptor ligands and probes derived from this sequence can be
CC used to assay for GABA-B receptors or DNA encoding them.
SQ Sequence 2924 BP; 628 A; 852 C; 793 G; 651 T;

Query Match 9.8% Score 317.2; DB 1; Length 2924;
Best Local Similarity 48.2%; Pred. No. 3.3e-68;
Matches 1034; Conservative 0; Mismatches 1083; Indels 27; Gaps 4;

Qy 232 CCCCCGGCGCCATGCCAGTTGCCCGCGGCTCTGTAGCGGCCGCTCTCCATCATG 291
Db ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
Qy 271 CTCGGCGGCTCACTCGGGTCCCGCCGACCCCTCCAGAACGGCGGCGAGTAC 330
Db ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
Qy 292 GGCCTCATGCGCTCACCAGAGGTTGGCCAGGAGGAGCATCGGGCGGGTGTGCTCCC 351
Db ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
Qy 331 ATCGGGGCACTGTTTCCCATGAGCGGGGGTGGCCAGGGGCCAGGCTGCCAGCGCG 390
Qy 352 GCCGTGGAACCTGGCATCGAGCAGATCCGGAACGAGTCACTCTGCGCCCTACTTCTC 411
Db ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
Qy 391 GTGAGATGGCGCTGGAGGACGTGAATAGCGGAGGACATCTCTCCGGACTATGAGCTC 450
Qy 412 GACCTGCGGCTTATGACGAGGTGGCAACAGCGAAAGGTTGAAGCCTTCTACGAT 471
Db ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
Qy 451 AAGCTCATCCACGACAGCAAGTGTGATCCAGGCCAACCCACCAAGTACCTATGAG 510
Db ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
Qy 472 CGGATAAATACGGCGGACCACTTGTATGGTGTGTTGGAGCGGTCTGTCCATCGGTACA 531
Db ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
Qy 511 CTGCTTACAACGACCTATCAA---GATCATCTTATGCTGGCTGCAGCTCTGTCTCC 567
Db ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||

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|----|------|------|--|------|
| Qy | 1606 | GC | TTTTCCTCTTCTTCAACATCAAGAACCGGAATCAGAAGCTCATAAAGATGTCGAGTCCA | 1665 |
| Db | 1639 | GT | CTGTCTGTCTCTTTTAACTATACAACTACACATCAGATGCTCCCTTATATCCGAAGATCACAGCCC | 1698 |
| Qy | 1666 | TAC | ATGAACAACCTTATCATCTCTGGAGGGATGCTTTTCCCTATGTCATCCATATTTCTCTTT | 1725 |
| Db | 1699 | AAC | CTGAACAACTGACTGCTGTGGCTGCTCAGCTGGCTTTAGCTGTGTCCTCCCTCG | 1758 |
| Qy | 1726 | GG | CTTGATGGATCCCTTTTCTCTGTAAGAACCTTTGAAACACTTTGCACGTCGAGACC | 1785 |
| Db | 1759 | GG | CTCGATGGTTACCACATTTGGAGGAACAGGATTTCTCTGCTGCGAGCGCGCTC | 1818 |
| Qy | 1786 | TGA | TTCTACACGTGGGCTACAGCACCGCTTTTGGGGCCATGTTTGGAAAGCACTCGAGA | 1845 |
| Db | 1819 | TGG | TCTCTGGGCTTGA | 1878 |
| Qy | 1846 | GT | CCAGCCATCTTCAAAATGTGAATAGAAAG- - - - -GAGATCATCAAGGAC | 1896 |
| Db | 1879 | GT | CCACAGCTCTTCAAGAAGGAAGAAAGAGGAGTGGAGGAAGACTCTGGAACCC | 1938 |
| Qy | 1897 | CAG | AAACTGCTTGTGATCGTGGGGGCATCGTCGTGATCGACACTGTGTATCCTGATCTGC | 1956 |
| Db | 1939 | TGG | AAGCTGTATGCCACGTGGGCTGTGGTGGCGATGCTCTCACTCTCGCCATC | 1998 |
| Qy | 1957 | TGG | CAGGCTGTGAGCCCTCGAAGGACAGTGGAGAGATACAGCATGGAGCCGAGACCCA | 2016 |
| Db | 1999 | TGG | CAGATCGTGGACCTCTGCACCGGACATTGAGACATTTGCCAAGGGAACCTAAG | 2058 |
| Qy | 2017 | GC | AGGAGGATATCTCCATCCGCTCTCTCTGGAGCACTGTGAGAAACCCCATATGACC | 2076 |
| Db | 2059 | GA | AGATATTACGCTCTCTATTCTGCCCCAGCTTGGAGCATTTGCAGTCCAGGAAGATGAAT | 2118 |
| Qy | 2077 | AT | CTGGCTTGGCATCTCTATGCTACAAAGGACATCTCATGTTTTCGGTGTGTTTCTTA | 2136 |
| Db | 2119 | ACA | TGGCTTGGCATTTCTATGTTACAGGGGCTGCTGCTGCTCTGSGAATCTCTCT | 2178 |
| Qy | 2137 | GCT | TGGAGACCCGCAACGTCAGCATCCCGCACTCAACGACAGCAAGATACATCGGATG | 2196 |
| Db | 2179 | GCT | TATGAGACCAAGATGTGTCCACTGAGAAGATCAATGATCACCGGCTGTGGGATG | 2238 |
| Qy | 2197 | AG | TCTTACACGCTGGGATCAGTGCATCAGCGGGCGCTGTCTCTCTGACCGG | 2256 |
| Db | 2239 | GCT | ATCTACAAATGTGGCAGTCTCTGTGCCCTCATCTGCTCTGTCAACATGATCTGTCC | 2298 |
| Qy | 2257 | GACC | AGCCCAATGTGCAGTTCTGCATCTGTGGCTCTGGTTCATCATCTTCTTGACGACCACT | 2316 |
| Db | 2299 | AG | CAGCAGGATCGACCTTTGCCCTTGGCTCTCTTGCCATAGTTTCTCTCTCTATATC | 2358 |
| Qy | 2317 | ACC | CTCTGCTGTATTCTGTGCGGAAGCTCATACCCCTGAGAAC | 2360 |
| Db | 2359 | ACT | CTGTGTGTCTTTGTGCCAAGATGCGAGGCTGATCAC | 2402 |

| | | |
|--------|---|-----------------------|
| RESULT | 2 | |
| VI0265 | | |
| ID | VI0265 standard; cDNA to mRNA; 2620 BP. | |
| AC | VI0265; | |
| DT | 03-JUN-1998 (first entry) | |
| DE | Human GABA-BR1a/b receptor cDNA. | |
| KW | Gamma-aminobutyric acid; GABA-BR1a/b receptor; human; brain; agonist; | |
| KW | inhibitory neurotransmitter; peripheral nervous system; antagonist; | |
| KW | treatment; dementia; depression; anxiety; bronchial inflammation; asthma; | |
| KW | epilepsy; cognitive function; ds. | |
| OS | Homo sapiens. | |
| FH | Key | Location/Qualifiers |
| FT | CDS | 1..2382 |
| FT | | /*tag= a |
| FT | | /product= GABA-BR1a/b |
| FT | | |
| PN | W03746675-A1. | |
| PD | 11-DEC-1997. | |
| PF | 19-MAR-1997; E01370. | |
| PR | 22-NOV-1996; US-756091. | |
| PR | 30-MAY-1996; US-655716. | |

QY 2191 GGGATGAGTGTCTACAACTGGGGATCATGTGCATCATCGGGCGCGTGTCTCTCTCTG 2250
 Db 1912 GGCATGGCTATCTACAATGTGGCAGTCTGTGGCTCATCACTGCTCTCTCTCTCTG 1971
 QY 2251 ACCCGGGACACCAATGTGAGTTCATGATGCTGTGGCTGTGGCTCATCATCTCTG 2310
 Db 1972 CTGTCAGCAGCAGGATGAGCTTTGCTCTCTCTCTCTCTCTCTCTCTCTCTCTCC 2031
 QY 2311 ACCATCACCTCTGCTGTGATTCGCGGAGCTCATCACTCATCACTCATCACTCAT 2360
 Db 2032 TATATCACTCTGTTGTGTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 2081

RESULT 3
 V10266
 ID V10266 standard; cDNA to mRNA; 2837 BP.
 AC V10266;
 DE 03-JUN-1998 (first entry)
 DE Rat GABA-BR1b receptor cDNA.
 KW Gamma-aminobutyric acid; GABA-BR1b receptor; rat; brain; agonist;
 KW inhibitory neurotransmitter; peripheral nervous system; antagonist;
 KW treatment; dementia; depression; anxiety; bronchial inflammation; asthma;
 KW epilepsy; cognitive function; ds.
 OS Rattus norvegicus.
 FH Key Location/Qualifiers
 FT CDS 228..2762
 FT /*tag= a
 FT /product= GABA-BR1b
 FN W09746675-A1.
 PD 11-DEC-1997.
 PF 19-MAR-1997; E01370.
 PR 22-NOV-1996; US-756091.
 PR 30-MAY-1996; US-655716.
 PA (NOVS) NOVARTIS AG.
 PI Bettler B, Bittiger H, Froestl W, Kaupmann K, Mickel SJ;
 PI WPI; 98-042183/04.
 DR P-PSDB; W40118.
 PT Purified GABA-B receptor or receptor protein - and antagonists of
 PT these which may be useful in treating nervous system disorders
 PS Claim 3; Page 67-74; 108pp; English.
 CC This cDNA sequence encodes a novel rat GABA-B receptor protein,
 CC GABA-BR1b. GABA (gamma-aminobutyric acid) is the major inhibitory
 CC neurotransmitter found in the brain and peripheral nervous system
 CC and this receptor may be used for the identification of GABA-B
 CC receptor agonists and antagonists. Such proteins may be used in
 CC treatment of dementia, depression, anxiety, epilepsy, spasticity,
 CC bronchial inflammation or asthma or to improve cognitive function.
 CC GABA-B receptor ligands and probes derived from this sequence can be
 CC used to assay for GABA-B receptors or DNA encoding them.
 SQ Sequence 2837 BP; 621 A; 842 C; 764 G; 610 T;

Query Match 9.6%; Score 312.8; DB 1; Length 2837;
 Best Local Similarity 48.0%; Pred. No. 3.9e-67;
 Matches 1038; Conservative 0; Mismatches 1097; Indels 27; Gaps 4;

QY 214 GCGGCTCCGCGCTGACACCCCGCGGCGCATGCCCAGTTGCCCGCGCTCTGTAG 273
 Db 312 GCCTCTCACTCCCTCATCTCCCGCGGCTCACCGAGGTTCCCGCGCACCCCTCTCA 371
 QY 274 GGCCGCTCTCCATCATGCGCTCATGCGCTCACCAAGAGTGCGCAAGGCGAGCATC 333
 Db 372 GAACGGCGCTGAGTATACATCGGGGCGTGTTCCTCCATGAGCGGGGCTGGCGGGGC 431
 QY 334 GGGCGGCTGCTCCCGCGTGGAACTGCGCATCGACAGATCCGCAACGATCATCTC 393
 Db 432 CAGCGCTCCGACCGCGGCTGGAGATGGCGCTGGAGAGCGTTAACAGCGCAGAGCATC 491
 QY 394 CTGCGCCCTCTCTCTCTGACCTCGCGCTCTATGACACGAGTGCACCAACGCAAGGG 453
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QY 454 TTGAAGCCCTTACAGTCGGATATAAATACGGGCGAACCACCTTGTGGTGTTCGAGGC 513
 Db 552 ACCAAGTACTTGTGACAACTACTTACAATGATGCCCCATCAAGATCATTTCTATGCTGG- 611
 QY 514 GTCTGTCCATCGGTACATCCATCATTTGAGAGTCCCTCAAGGCTGGAATCTGTGTGAG 573
 Db 611 --CTGTAGTCTGTCTCCACACTTGTAGCTGAGGCTGCCCGATGTGNAACCTTATGTG 568
 QY 574 CTCTCTCTCTCTGCAACACCGCTGTTCAGCGGATTAAGAAAAATACCTTTATTTCTTT 633
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 QY 994 CACACGGAAGCAACTCATCCCGCTCCGCGGAAAGAACTGCTTCTGCTCCATCGAGGCG 1053
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 QY 1054 TACATTGGCTGGATTTGAGCGCCCTGAGCTCCAAGCAGATCAAGACCATCTCAGAGAA 1113
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 QY 1114 ACTCCACAGCATGATGAGAGAGATACAAACAACAGCGTTCAG-----CGTGGGCGCC 1167
 Db 1200 ACGTCACAGGAATTTGTGGAGAAACTAACCAAGCGCTGAAAAGACACCCCGAGAGACT 1259
 QY 1168 AGCAAGTTCACGGGTACGCTAGATGGCATCTGGGTATCCCAAGACACTGACAGAGS 1227
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Db 1620 AAGACATCCGTTCTCTCAGAAACTCTTTATCTCCGTCAGTTCTCTCCAGCGCTG 1679
QY 1588 GGGATGATCATGGCCAGTCTTTCTCTTCAACATCAAGAACCGGAATCAGAGCTC 1647
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Db 1860 GTCTGCCAGCCCGCTTTGGCTCTTGGGCTTGGGCTTTAGTCTGGGCTATGGCTCTATG 1919
QY 1828 TTTGCAAGACCTGGAGTCCACGCCATCTTCAAAAATGTGAAATGAAGAA----- 1881
Db 1920 TTCACCAAGATCTGTGGTCCACACAGCTTTCAGGAGAGGAGGAGAGAGGAGTGG 1979
QY 1881 --GAGATCATCAAGACACAGAACTGTTGTGATCGTGGGGGATGCTCTGATCGAC 1938
Db 1980 AGGAAGACCCCTAGAGCCCTGGAAACTCTATGCCACTGTGGCCCTGCTGTGGCATGGAT 2039
QY 1939 CTGTGTATCTCTGATCTGTGGCAGCTGTGGACCCCTCGGACGACAGTGGAGAGTAC 1998
Db 2040 GTCTCGACTCTTGGCATCTGGCAGATTGTGGACCCCTTGGACCCGATTTGAGACATTTT 2099
QY 1999 AGCATGAGCCGGACCCAGCAGGAGGGATATCTCCATCCGCCCTCTCTCTGGAGCACTGT 2058
Db 2100 GCCAAGGAGGAACCAAGAGACATCGATGCTCTCATTTGCCAGTTGGACCACTGC 2159
QY 2059 GAGACACCCATATGACCATCTGGCTTGGCATCGTGTATGCTTACAGGGACTTCTCATG 2118
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QY 2359 AC 2360
Db 2460 AC 2461

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RESULT 4

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V10264
ID V10264 standard; cDNA to mRNA; 4376 BP.
AC V10264;
DT 03-JUN-1998 (first entry)
DE Rat GABA-BR1a receptor cDNA.
KW Gamma-aminobutyric acid; GABA-BR1a receptor; rat; brain; agonist;
KW inhibitory neurotransmitter; peripheral nervous system; antagonist;
KW treatment; dementia; depression; anxiety; bronchial inflammation; asthma;
KW epilepsy; cognitive function; ds.
OS Rattus norvegicus.
FH key Location/Qualifiers
FT CDS 182..3064

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FT
FT
PN
PD
PF
PR
PR
PA
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PI
DR
DR
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PS
CC
CC
CC
CC
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CC
SQ
Query Match
Best Local Similarity
Matches 967; Conservative 0; Mismatches 990; Indels 27; Gaps 4;
QY 392 TCCTGCGCCCTCTACTTCTCTGACCTCGGCTCTATGACACGGAGTGGCAGACGCAAAAG 451
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Db 1269 GACTTTTCTATGAGCGGAAGCCCGGAAAGTTTGTGTGAGGCTCTATAAGAAAGGCTCT 1328
QY 932 ATGGTAGTAAATATCAGTGGATCATTCGCGGCTGGTACGAGCTTCTTGGTGGGAGCAGG 991
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WO9746675-A1.
11-DEC-1997.
19-MAR-1997. E01370.
22-NOV-1996; US-756091.
30-MAY-1996; US-655716.
(NOV5) NOVARTIS AG.
PI Bettler B, Bittiger H, Froestl W, Kaupmann K, Mickel SJ;
WPI: 98-042183/04.
P-PSDB: W40116.

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PT Purified GABA-B receptor or receptor protein - and antagonists of
PT these which may be useful in treating nervous system disorders
PS Claim 3: Page 42-50; 108pp; English.
CC This cDNA sequence encodes a novel rat GABA-B receptor protein,
CC GABA-BR1a. GABA (gamma-aminobutyric acid) is the major inhibitory
CC neurotransmitter found in the brain and peripheral nervous system
CC and this receptor may be used for the identification of GABA-B
CC receptor agonists and antagonists. Such proteins may be used in
CC treatment of dementia, depression, anxiety, epilepsy, spasticity,
CC bronchial inflammation or asthma or to improve cognitive function.
CC GABA-B receptor ligands and probes derived from this sequence can be
CC used to assay for GABA-B receptors or DNA encoding them.
SQ Sequence 4376 BP; 940 A; 1209 C; 1195 G; 1032 T;

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QY 1052 GCTACATTGGCGTGGATTTCAGAGCCCTGAGCTCCAAAGCAGATCAAGACCACTCTCAGGAA 1111
 Db 1440 GCCACATCACCGAGGATGCTCATGCTGACCTGCCACACCCGAGCATTTCCAAACA 1499
 QY 1112 AGACTCCACAGCATGATGAGAGAGAGTACAAACAAGCGGTACG-----CGTGGGGC 1165
 Db 1500 TGACGTCACGAATTTGTGAGAGAACTTAACCAAGCGCTGAAAGACACCCCGAGGAGA 1559
 QY 1166 CCAGCACTTCCAGCGTACCGCTACGATGCATCGGCTCATCGCCCAAGACACTGCAGA 1225
 Db 1560 CTGGAGGCTTCCAGAGGACCACTGGCCTATGATGCTATCTGGGCCCTTGGCTTTGGCCT 1619
 QY 1226 GGGCCATGGAGACACTGATGCCAGCAGCGCCACAGCGGATCCAGGACTTCAACTACA 1285
 Db 1620 TGAACAAGACGCTGGAGGAGTGTGCTTCCGCGCTGGCCTGGAGGACTTTAACTACA 1679
 QY 1286 CGGACCAACGCTGGCGAGGATCATCTCAATGCCATGAAGACCAACACTTCTTCGGGG 1345
 Db 1680 ACAACGACGACATTACAGACGAGATACCGGGCCATGAACCTCCTCTCTTTGAGGGCG 1739
 QY 1346 TCACGGGTCAAGTTGATTTCCGGAATGGGAGAGAATGGGACCATTAATTTACTCAAT 1405
 Db 1740 TTTCTGGCCATGTGCTTTGATGCGAGCGCTCCCGATGGCATGGACACTTATCGAGC 1799
 QY 1406 TTCAGACAGCAGGAGGTGAAGTGGGAGAGTAAACGCTGTGGCGGACACACTGGAGA 1465
 Db 1800 AGCTACAGGCGGAGCTACAAAGAGATCGGCTACTAGCAGACCAACGAGTATGATCTT 1859
 QY 1466 TCATCAATGACACCATCATAGTTTCAAGATCCGAACCAACCAAGACAAAGACCAATCC 1525
 Db 1860 CTTGTCCAAAACGACAAAGTGGATGGAGGGTCTCCCGCAGCTGACGACACTTGTGTA 1919
 QY 1526 TGGACGAGCTCGGAAGATCTCCCTACCTCTACAGCATCTCTCTGCGCTCACCAATCC 1585
 Db 1920 TCAAGACATTCGTTCTGCTCTCAGAACTCTTTATCTCGGCTCAGTCTCTCCAGCC 1979
 QY 1586 TCGGATGATCATGCCAGTCTTTTCTCTTCTCAACATCAAGAACCGGAATCAGAACG 1645
 Db 1980 TGGGATGTTCTGCTGCTGCTGCTCTTAACTATCAACTCCACACTCCACGCTTGGT 2039
 QY 1646 TCATAAAGATGCGAGTCCATACATGAACACCTTATCACTTGGAGGATGCTTCCCT 1705
 Db 2040 ATATCCAGATCCCGACCCCACTGACATCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 2099
 QY 1706 ATGCTTCCATATTTCTTTTGGCTTGTATGATGCTTTGCTCTGAAAGACCTTTGAAA 1765
 Db 2100 TGGCTGCTCTTCTCTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 2159
 QY 1766 CACTTTGACCGCTCAGGACCTTGTACCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1825
 Db 2160 TTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 2219
 QY 1826 TGTGTTCAAGACCTGAGAGTCCAGCCATCTTCAAAATGTGAAATGAAGAA----- 1881
 Db 2220 TGTTCACCAAGATCTGTGGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 2279
 QY 1881 ----GAAGATCATCAAGACCAAGAACTGCTGTGATGCTGGGGGCTGCTGCTGCTGCT 1936
 Db 2280 GGAGGAAGACCTTAGAGCCCTGGAACCTATATGCTGCTGCTGCTGCTGCTGCTGCTGCT 2339
 QY 1937 ACCTGTGATCCTGATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1996
 Db 2340 ATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 2399
 QY 1997 ACAGCATGGAGCGGACCCAGCAGGAGGATATCTCATCCGCCCTCTCTCTGGAGCACT 2056
 Db 2400 TTGCAAGGAGGAAACCAAGGAAGACATCGATGTCTCCATCTCTCCCGCAGTGGAGCACT 2459
 QY 2057 GTGAGAACACCATATGACCATCTGGCTTGGCATCGTCTATGCTTACAGGAGGACTTCTCA 2116
 Db 2460 GCAGCTCCAAAGATGATACGTGGCTTGGCATTTCTATGTTTACAGGAGGCTGCTGC 2519

QY 2117 TGTGTTCCGTTGTTTACGTTGGGAGACCCGCAACAGTCAGCATCCCGCACTCAAGC 2176
 Db 2520 TGTCTCTGGGAATCTTTCTGTTACGAAACCAAGAGCGTCTCCACTGAAAGATCAATG 2579
 QY 2177 ACAGCAAGTACATCGGATGAGTGTCTACACGCTGGGATCATGTGCATCATCGGGCGC 2236
 Db 2580 ACCACAGGCGCTGGGCTATCTACAAATGTCGCGTCTGCTGCTCATCACTGCTC 2639
 QY 2237 CTGCTCTCTCTGACCCCGGACGACCAATGTGCACTTCTGATGCTGCTGCTGCTGCT 2296
 Db 2640 CHTGACCATGATCTTCCAGTCAGCAGGACGAGCCTTTGGCTTTGCTCTCTGGCCA 2699
 QY 2297 TCATCTTCTGACGACCATCCCTCTGCTGTTGTTGTCGCCGAAGCTCATCACTCCGTA 2356
 Db 2700 TCGTGTCTCTCTTACATCACTCTGCTGTTGCTCTTTGTCGCCAAGATGGCAGGCTGA 2759
 QY 2357 GAAC 2360
 Db 2760 TCAC 2763

RESULT 5
 V88919
 ID V88919 standard; cDNA; 314 BP.
 AC V88919;
 DT 12-FEB-1999 (first entry)
 DE EST clone HW456.
 KW Expressed sequence tag; secreted protein; haematopoiesis regulator;
 KW tissue growth; activin; inhibin; tumour invasion suppressor; EST; human;
 KW chemotaxis; chemokinesis; haemostasis; gene therapy; thrombolysis;
 KW receptor; ligand; anti-inflammatory; tumour inhibitor; ds.
 OS Homo sapiens.
 PN W09845437-A2.
 PD 15-OCT-1998.
 PE 10-APR-1998; U06956.
 PR 10-APR-1997; US-837312.
 PA (GENY) GENETICS INST INC.
 PI Agostino MJ, Jacobs K, Lavallie ER, McCoy JM, Merberg D,
 PI Racie LA, Spaulding V, Treacy M;
 DR WPI: 99-070078/06.
 PT New polynucleotides encoding human secreted proteins - derived from
 PT e.g. human blood, kidney, foetal lung, placenta, testes, brain,
 PT ovary, pituitary, retina and colon cDNA libraries
 PS Claim 1; Page 556-557; 641pp; English.
 CC The present sequence represents an expressed sequence tag (EST), and is
 CC a polynucleotide of the invention. The polynucleotides of the invention
 CC are all secreted EST sequences isolated from a variety of human tissue
 CC sources. The EST sequences and proteins encoded by them are predicted to
 CC have useful biological activities which would make them suitable for
 CC treating, preventing or ameliorating medical conditions in humans and
 CC animals, although no supporting data is given. Suggested activities
 CC include nutritional activity, immune stimulating or suppressing activity,
 CC haematopoiesis regulating activity, tissue growth activity,
 CC activin/inhibin activity, chemotactic/chemokinetic activity, haemostatic
 CC and thrombolytic activity, receptor/ligand activity, anti-inflammatory
 CC activity, cadherin/tumour invasion suppressor activity, tumour inhibition
 CC activity. The EST sequences are also stated to be useful for gene
 CC therapy.
 SQ Sequence 314 BP; 72 A; 81 C; 89 G; 72 T;

Query Match 8.8%; Score 286.8; DB 1; Length 314;
 Best Local Similarity 97.7%; Pred. No. 3.6e-61;
 Matches 291; Conservative 0; Mismatches 7; Indels 0; Gaps 0;

QY 1857 CTTCAAAATGTGAAATCAAGACAGATCATCAAGCAGCAACTGCTGTGATCGT 1916
 Db 10 CTTTCATGCGCTGAAATGAAGAGAAATCATCAAGGACCAAGCAACTGCTGTGATCGT 69
 QY 1917 GGGGGGCGATGCTGCTGATCGACCTGTGATCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1976
 Db 70 GGGGGGCGATGCTGCTGATCGACCTGTGATCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 129

QY 1977 GCGAAGACAGTGGAGAGTACAGCATGGAGCCGGACCCAGCAGGAGGATATCTCCAT 2036
 Db 130 GCGAAGGACAGTGGAGAGTACAGCATGGAGCCGGACCCAGCAGGAGGATATCTCCAT 189
 QY 2037 CGCGCCCTCTCTGGAGCACTGTGAGAACACCCATATGACCATCTGGCTTGGCATCTCTA 2096
 Db 190 CGCGCCCTCTCTGGAGCACTGTGAGAACACCCATATGACCATCTGGCTTGGCATCTCTA 249
 QY 2097 TGCTTACAGGACATCTCATGTGTGTTGGTGTGTTCTTAGCTTGGGAGACCCGCAAC 2154
 Db 250 TGCTTACAGGACATCTCATGTGTGTTGGTGTGTTCTTAGCTTGGGAGACCCGCAAC 307
 RESULT 6
 X51891
 ID X51891 standard; DNA: 361 BP.
 AC X51891.
 DT 22-JUN-1999 (first entry)
 DE Human secreted protein 5', EST SEQ ID NO: 105.
 KW Human; secreted protein; EST; expressed sequence tag; diagnosis;
 KW forensic; gene therapy; chromosome mapping; signal peptide;
 KW upstream regulatory sequence; cytokine activity; cell proliferation;
 KW differentiation; haematopoiesis regulation; tissue growth regulation;
 KW reproductive hormone regulation; chemotactic; chemokinetic; haemostatic;
 KW thrombolytic; anti-inflammatory; tumour inhibition; ds.
 OS Homo sapiens.
 PN WO9906552-A2.
 PD 11-FEB-1999.
 PF 31-JUL-1998; IB1236.
 PR 01-AUG-1997; US-905223.
 PA (G8ST) GENSET.
 PI Duclert A, Dumas Milne Edwards J, Lacroix B;
 DR WPI; 99-153782/13.
 DR P-PSDB; Y13091.
 PT New isolated brain-derived nucleic acids - used to develop products
 PT which may have cytokine, immune, regulatory, haematopoiesis
 PT regulating, anti-inflammatory or tumour inhibition activity
 PS Claim 1; Page 253; 57pp; English.
 CC X51787 to X52019 represent 5' expressed sequence tags (ESTs) for human
 CC secreted proteins, and encode the proteins given in Y12987 to Y13219,
 CC respectively. The proteins given represent the signal peptide and an
 CC N-terminal fragment of a secreted protein. The nucleic acid sequences
 CC can be used for producing secreted human gene products. They can also
 CC be used to develop products for diagnosis and therapy. The proteins
 CC obtained may have cytokine activity, cell proliferation/differentiation
 CC activity, haematopoiesis regulating activity, tissue growth regulating
 CC activity, reproductive hormone regulating activity, chemotactic/
 CC chemokinetic activity, haemostatic and thrombolytic activity, receptor/
 CC ligand activity, anti-inflammatory activity, tumour inhibition activity
 CC or other activities. The products can be used in forensic, gene therapy
 CC and chromosome mapping procedures. The sequences can also be used for
 CC obtaining corresponding promoter sequences. The nucleic acids encoding
 CC the signal peptide can be used for directing extracellular secretion of
 CC a polypeptide or the insertion of a polypeptide into a membrane, or
 CC importing a polypeptide into a cell.
 SQ Sequence 361 BP; 86 A; 89 C; 94 G; 92 T;
 Query Match 5.58; Score 177; DB 1; Length 361;
 Best Local Similarity 90.48; Pred. No. 2.9e-34;
 Matches 189; Conservative 0; Mismatches 20; Indels 0; Gaps 0;
 QY 404 ACTTCTCTCGACCTGGCGTCTATGACACGGAGTGGACAGCAACGAAAGGGTTGAAGCCT 463
 Db 153 AGTCATCTATTGGAGCGGCAATCATCTCAGTGGGACACGAAAGGGTTGAAGCCT 212
 QY 464 TCTAGATGCGATAAATACGGGGCGAACCACTTGATGGTGTGTTGGAGGCGTCTGTCCAT 523
 Db 213 TCTAGATGCGATAAATACGGGGCGAACCACTTGATGGTGTGTTGGAGGCGTCTGTCCAT 272
 QY 524 CCGTCACATCATCATTCGACAGTCCCTCCAGGCTGGAATCTGGTGCAGGTTTCTTTG 583
 Db 273 CCGTCACATCATCATTCGACAGTCCCTCCAGGCTGGAATCTGGTGCAGGTTTCTTTG 332

QY 584 CTGCAACCAACGCTGTCTTAGCCGATAAG 612
 Db 333 CTGCAACCAACGCTGTCTTAGCCGATAAG 361
 RESULT 7
 Q11726
 ID Q11726 standard; DNA: 2176 BP.
 AC Q11726.
 DT 25-JUN-1991 (first entry)
 DE Sequence encoding viral reverse transcriptase enzyme.
 KW Reverse transcriptase; HIV; HTLV-I; ss.
 OS Myxococcus xanthus.
 FH Key Location/Qualifiers
 FT cds 640..2097
 FT /*tag= a
 FT /product= viral reverse transcriptase
 PN J03022975-A.
 PD 31-JAN-1991.
 PF 22-FEB-1990; 042305.
 PR 24-FEB-1989; US-315316.
 PA (UYNE-) Univ of New Jersey.
 DR WPI; 91-127415/18.
 DR P-PSDB; R11919.
 PT Novel protein having reverse transcriptase enzyme activity -
 PT obtd. by culturing Myxococcus xanthus FB
 PS Disclosure; fig 4; 9pp; Japanese.
 CC This sequence encodes a protein with viral (e.g. HIV or HTLV-I)
 CC reverse transcriptase enzyme activity and is extracted from
 CC Myxococcus xanthus. See also J03022976.
 SQ Sequence 2176 BP; 340 A; 763 C; 789 G; 284 T;
 Query Match 1.48; Score 45.4; DB 1; Length 2176;
 Best Local Similarity 50.68; Pred. No. 0.1;
 Matches 135; Conservative 0; Mismatches 131; Indels 1; Gaps 1;
 QY 149 GGAGTCGAGGGCGGAGGAGAGCCGCTGAGTGAGCAGAGTCACAGCGCTGCGCCCCA 208
 Db 891 GAAGGCTTGAGAGGAGAGAGAGCCGAGGCGCCGCCCTGAAGCGTCA 950
 QY 209 GAATCGCGCTCCGCCCTGACACCCCGCGCCATGCCCCAGTTGCCCGCGCTCTG 268
 Db 951 GCGCACAGGGCTGGAAGCCACGACGTGGCCACCTGCGCCGCCCTG 1009
 QY 269 CTAGGGCGCGCTCTCCATCATGCGGCTCATCGCGTACCAAGAGGTTGGCAAGGCA 328
 Db 1010 CGAGAGACCCCTGCGCGACGGGTTGAGTGTGCGCCACCGCGAGGAGCGCCCGGCA 1069
 QY 329 GCATCGGGCGGCTGTGCTCCCGCGGTGAACTGCGCATCGAGCAGATCCGCAACGAGT 388
 Db 1070 ACGGCTGAGGAGCTGGAATCGGGGAGGCGCTGCGCAAGGCGCTGGGCTGACCGTGT 1129
 QY 389 CACTCTCGCGCCCTACTTCTCTCGACC 415
 Db 1130 CCAAGCTCGCTGTTCTGCTCCACC 1156
 RESULT 8
 X51965
 ID X51965 standard; DNA: 397 BP.
 AC X51965.
 DT 22-JUN-1999 (first entry)
 DE Human secreted protein 5', EST SEQ ID NO: 179.
 KW Human; secreted protein; EST; expressed sequence tag; diagnosis;
 KW forensic; gene therapy; chromosome mapping; signal peptide;
 KW upstream regulatory sequence; cytokine activity; cell proliferation;
 KW differentiation; haematopoiesis regulation; tissue growth regulation;
 KW reproductive hormone regulation; chemotactic; chemokinetic; haemostatic;
 KW thrombolytic; anti-inflammatory; tumour inhibition; ds.
 OS Homo sapiens.
 PN WO9906552-A2.

11-FEB-1999. PF
31-JUL-1998; IB1236. PF
01-AUG-1997; US-905223. PR
(GENT) GENSET. PR
PI
PI Duclert A, Dumas Milne Edwards J, Lacroix B; DR
WPI: 99-153782/13. DR
P-PSDB; Y13165. DR
PT
PT New isolated brain-derived nucleic acids - used to develop products
PT which may have cytokine, immune, regulatory, haematopoiesis
PT regulating, anti-inflammatory or tumour inhibition activity
PT
PT Claim 1: Page 349; 577pp; English. PT
PS
CC X51787 to X52019 represent 5' expressed sequence tags (ESTs) for human
CC secreted proteins, and encode the proteins given in Y12987 to Y13219,
CC respectively. The proteins given represent the signal peptide and an
CC N-terminal fragment of a secreted protein. The nucleic acid sequences
CC can be used for producing secreted human gene products. They can also
CC be used to develop products for diagnosis and therapy. The proteins
CC obtained may have cytokine activity, cell proliferation/differentiation
CC activity, haematopoiesis regulating activity, tissue growth regulating
CC activity, reproductive hormone regulating activity, chemotactic/
CC chemokinetic activity, homeostatic and thrombolytic activity, receptor/
CC ligand activity, anti-inflammatory activity, tumour inhibition activity
CC or other activities. The products can be used in forensic, gene therapy
CC and chromosome mapping procedures. The sequences can also be used for
CC obtaining corresponding promoter sequences. The nucleic acids encoding
CC the signal peptide can be used for directing extracellular secretion of
CC a polypeptide or the insertion of a polypeptide into a membrane, or
CC importing a polypeptide into a cell.
SQ Sequence 397 BP; 79 A; 111 C; 93 G; 107 T;

Query Match 1.4%; Score 45.4; DB 1; Length 397;
Best Local Similarity 51.3%; Pred. No. 0.05;
Matches 117; Conservative 3; Mismatches 107; Indels

| | | | |
|----|------|--|------|
| Qy | 1501 | CCACCAAAAGACAAAGACCATCATCTGGAGACAGCTGCCGAGAGATCTCCCTACCTCTCTAC | 1560 |
| | | | |
| Db | 168 | CCCCCAGCTGSRGAGACCCCTGGTCTCATCAAGACATTCGCGTCTCTGTCTAC-ASANNCTTTT | 226 |
| | | | |
| Qy | 1561 | AGCATGCTCTCTGCCCTCACCATCTCGGGATGATCATGGCCAGTGCTTTTCTCTCTTC | 1620 |
| | | | |
| Db | 227 | ATCTCGCTCTCAGTTCTCTCCAGCTGGGCATGTCTAGCTGTTGTCTGCTCTCTTT | 286 |
| | | | |
| Qy | 1621 | AACATCAAGACCGGAATCAGAGCTCATAAAGATGTGAGTCCCATCATGAACAACGTT | 1680 |
| | | | |
| Db | 287 | AACATCTACAACCTCACATGCTCGGTATATCCAGAACTCACAGCCCACTGAAACAACCTG | 346 |
| | | | |
| Qy | 1681 | ATCATCTTGGAGGGANGTTTCCCTATGCTTCCATATTTCTTTGGC | 1728 |
| | | | |
| Db | 347 | ACTGCTGTGGGCTGTCAMTGGCTTTAGCTGCTTCTCCCTGGGGC | 394 |
| | | | |

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RESULT      9
Q10613/c
ID Q10613 standard; DNA; 15672 BP.
AC Q10613;
DT 26-APR-1991 (first entry)
DE Rianodin receptor gene.
KW Rianodin receptor gene; calcium release modulator; tranquiliser;
OS antagonist; ss.
OR Oryctolagus cuniculus.
FH Location/Qualifiers
FT 593..15553
FT /*tag= a
FT /*product= rianodin receptor
FT 351..356
FT /*tag= b
FT /*note= "feature unlabelled in specification"
FT 407..452
FT /*tag= c
FT /*note= "feature unlabelled in specification"
FT 392..396
FT /*tag= d
FT caat_signal
FT

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| | | | |
|----|---|--|---------|
| FT | | /note= "feature unlabelled in specification" | |
| FT | misc_feature | 503..510 | |
| FT | | /*tag= e | |
| FT | | /note= "feature unlabelled in specification" | |
| FT | misc_feature | 514..521 | |
| FT | | /*tag= f | |
| FT | | /note= "feature unlabelled in specification" | |
| FT | polya_signal | 15388..15393 | |
| FT | | /*tag= g | |
| FT | J03011098-A. | | |
| FT | 18-JAN-1991. | | |
| PD | 07-JUN-1989; | 144569. | |
| PR | 07-JUN-1989; | JP-144569. | |
| PA | (MITU) Mitsubishi Kasei Corp. | | |
| PA | WPI; 91-062003/09. | | |
| DR | P-PSDB; R10834. | | |
| DR | New rianodin receptor, genes encoding it and its prepn. - useful as | | |
| PT | calcium release modulator for tranquillisers and for assaying | | |
| PT | calcium antagonists. | | |
| PT | Disclosure; Fig 1; 18pp; Japanese. | | |
| CC | RNA confg. poly(A) was prepared from rabbit skeletal muscle | | |
| CC | endoplasmic reticulum. From the obtd. poly(A) mRNA, a cDNA bank | | |
| CC | corresp. to it was prepared by random primer method, Oligo (dT) | | |
| CC | primer method, and primer extension method. A cDNA was obtained by | | |
| CC | screening with a DNA probe (see Q10614-15). By introducing the obtd. | | |
| CC | cDNA into an expression vector, vector pRRS7 was formed. | | |
| CC | The product is said to be involved in calcium release from | | |
| CC | sarcoplasmic reticulum which triggers constriction of skeletal muscle. | | |
| CC | Therefore, the receptor is useful as tranquilliser and assay series for | | |
| CC | screening of calcium antagonist. | | |
| CC | Sequence 15672 BP; 3139 A. | | |
| SQ | Sequence 15672 BP; 3139 A. | 4909 C; 4955 G; | 2669 T; |

| Query Match | 1.3% | Score 43.6; | DB 1; | Length 15672; |
|-----------------------|---|-----------------|-----------|---------------|
| Best Local Similarity | 46.9%; | Pred. No. 0.63; | | |
| Matches 136; | Conservative 0; | Mismatches 154; | Indels 0; | Gaps 0; |
| Qy 2766 | TCCAGAACATCCAGCGTCGGCTGTCCCTCCAGCTCCCATCTCCACACAGCGCTACCT | 2825 | | |
| Db 13856 | TCGGGAGCCCTCGGGCGTGGCGGTTCCGCGCGTGGTGTGCCCCATGTGCCCAAGGC | 13797 | | |
| Qy 2826 | CCATCATCGGAGCGTGGAGCCACGCTGTTCAGGCCCTCGGTCAAGCCCCACCGCCAG | 2885 | | |
| Db 13796 | GGCAGCGCCCTCGGGCGCGGAAGGGGCCCCCGTGGGCCACGGCCACGACCCCTCGGCAC | 13737 | | |
| Qy 2886 | CCCCGGCCACAGACATGTGCCACCCCTCTTCCGAGTCATGCTCTCGGCGCCTGTAAAGGTG | 2945 | | |
| Db 13736 | CGCCCGGGCGGCGTCTGTGACCGCCACCTCCTCATCGCGTCACCCCTCGGCCGCGTCG | 13677 | | |
| Qy 2946 | GGAGCGTGGCGCGGGCGCTCCCGCTGACAGAACACACTGGGCAGAGGGGTCTGCTG | 3005 | | |
| Db 13676 | CCTCGCCTTCGCCCTTCGCCCGCGCGCGCTCCGCGTCCGCCCGCGGCCCGCGGCTGCTCGC | 13617 | | |
| Qy 3006 | CAGAAACACTGTCCGCTCTGGCTCGGAGAGCTGGGCACCACTATGGCTGGC | 3055 | | |
| Db 13616 | CGTGCACCTCTGCTGGTGGGATTCGGGCATGCCGGCCACGACGAGCTCGGCT | 13567 | | |

| | |
|--------|---|
| RESULT | 10 |
| V89458 | |
| ID | V89458 standard; cDNA; 273 BP. |
| AC | V89458; |
| DT | 15-FEB-1999 (first entry) |
| DE | EST clone CO337. |
| KW | Human; secreted protein; expressed sequence tag; EST; haematopoiesis; |
| KW | tissue growth; activin; inhibin; chemotaxis; chemokinesis; haemostatic; |
| KW | receptor; ligand; thrombolytic; anti-inflammatory; cadherin; anti-tumour; |
| KW | gene therapy; ss. |
| OS | Homo sapiens. |
| PN | W05845436-A2. |
| PD | 15-OCT-1998. |
| PF | 10-APR-1998; U06955. |
| PR | 10-APR-1997; US-838821. |

CC isolated from genomic DNA obtained from the human hepatoma cell line
 CC Hep 3B2. C/EBP's comprise a class of EBPs whose members are capable
 CC of preferentially recognising and binding a CCAAT sequence motif
 CC (such as is found in the transferin and ApoB genes), and enhancer
 CC core sequence motif, or the enhancer regions of several viral
 CC promoters. The DNA may be used in gene therapy to inhibit the
 CC proliferation of a tumour cell, esp. in antisense form, and may also
 CC be used to induce proliferation of a hepatic cell. Transgenic mice
 CC can be used to determine the carcinogenic potential of a chemical.
 CC Sequence 1312 BP; 197 A; 513 C; 471 G; 131 T;
 SQ

Query Match 1.3%; Score 41.8; DB 1; Length 1312;
 Best Local Similarity 54.1%; Pred. No. 0.63;
 Matches 85; Conservative 0; Mismatches 72; Indels 0; Gaps 0;

QY 2818 GCCTACTCCATCCATCGAGCGCTGGACGCCAGCTGTGTGAGCCCTCGTCAAGCCCC 2877
 DB 797 GCCAGACACCATGACCTGCAGCCGGTCAACCCAGCGCGCCGCGCCGCGCA 856
 QY 2878 ACCGCCAGCCCCGCCACACATGTGCGACCTCTCTCCGAGTATGCTCTCGGCGCTG 2937
 DB 857 GCCCGACCCCGCGCGCGCTCGTGGCGCGCGCTTCGGGCGCTGCGACGGCGCTCA 916
 QY 2938 TAAGGGTGGAGCGCTGGCGCGCGCGCTCCCGCGT 2974
 DB 917 AGGGCTGGCGCGCGCGCGCGCGCTCCCGCGAG 953

RESULT 14
 ID T44325 standard; DNA; 1312 BP.
 AC T44325;
 DT 27-JAN-1997 (first entry)
 DE DNA encoding CCAAT/Enhancer binding protein C/EBP-alpha.
 KW C/EBP-alpha; transcription factor; enhancer; cancer; treatment;
 KW gene therapy; CCAAT sequence motif; ss.
 OS Homo sapiens.
 FH Key Location/Qualifiers
 FT misc_feature 151..1233
 FT cds /tag= a
 FT /product= CCAAT/enhancer binding protein

US5545563-A.
 13-AUG-1996.
 PF 04-MAR-1993; 029325.
 PR 04-MAR-1993; US-029325.
 PR 04-MAR-1994; US-205506.
 PA (BAYU) BAYLOR COLLEGE MEDICINE.
 PI Darlington GJ, Wilde M, Willson DR;
 DR WPI: 96-383675/38.
 DR P-PSDB: W06797.
 PT DNA encoding human CCAAT-enhancer binding protein - useful for gene
 PT therapy of cancer, etc.
 PS Claim 2; Column 23-26; 23pp; English.
 CC The present sequence encodes human CCAAT/Enhancer Binding Protein (EBP)
 CC C/EBP-alpha (W06797). EBPs are transcriptional factors that bind to
 CC enhancer sequences. The ability of an enhancer to mediate gene expression
 CC in a particular cell is dependent upon the expression of an appropriate
 CC EBP in that cell. CCAAT/EBPs comprise a class of EBPs that are capable of
 CC preferentially recognising and binding a CCAAT sequence motif (such as
 CC found in the transferin and ApoB genes), an enhancer core sequence
 CC motif, or the enhancer regions of several viral promoters. The C/EBP-
 CC alpha DNA sequence is useful in gene therapy, esp. to treat cancer and
 CC other diseases. The present sequence differs from that described in
 CC U.S. patent application Ser. No. 8029325, most notably in including
 CC a 65 residue long NotI-NotI oligonucleotide (nucleotides 558-622 of the
 CC present sequence) between positions 556-557 of the prior sequence.
 CC Sequence 1312 BP; 197 A; 513 C; 471 G; 131 T;
 SQ

Query Match 1.3%; Score 41.8; DB 1; Length 1312;
 Best Local Similarity 54.1%; Pred. No. 0.63;
 Matches 85; Conservative 0; Mismatches 72; Indels 0; Gaps 0;

QY 2818 GCCTACTCCATCCATCGAGCGCTGGACGCCAGCTGTGTGAGCCCTCGTCAAGCCCC 2877
 DB 797 GCCAGACACCATGACCTGCAGCCGGTCAACCCAGCGCGCCGCGCGCA 856
 QY 2878 ACCGCCAGCCCCGCCACACATGTGCGACCTCTCTCCGAGTATGCTCTCGGCGCTG 2937
 DB 857 GCCCGACCCCGCGCGCGCTCGTGGCGCGCGCTTCGGGCGCTGCGACGGCGCTCA 916
 QY 2938 TAAGGGTGGAGCGCTGGCGCGCGCGCTCCCGCGT 2974
 DB 917 AGGGCTGGCGCGCGCGCGCGCGCTCCCGCGAG 953

RESULT 14
 ID 073500 standard; DNA; 8438 BP.
 AC Q73500;
 DT 15-MAY-1995 (first entry)
 DE DNA encoding Pseudorabies virus large latency transcript.
 KW Pseudorabies virus; PRV; LLV; large latency transcript;
 KW attenuated virus; vaccine; early protein 0; EP0; HSV-1 ICP0;
 KW protecting animals; deletion mutants; swine; ds.
 OS Pseudorabies virus.
 FH Key Location/Qualifiers
 FT misc_feature 1..7013
 FT /tag= a
 FT /note= "derived from PRV strain InPh"
 FT misc_feature 7014..8425
 FT /tag= b
 FT /note= "derived from PRV strain Ka"
 FT cds 622..6498
 FT /tag= c
 FT /note= "encodes predicted amino acid sequence of ORF2"
 FT tata_signal 1..6
 FT /tag= d
 FT misc_feature 34
 FT /tag= e
 FT /note= "RNA cap site"
 FT polya_signal 8382..8387
 FT /tag= f
 PN US5352596-A.
 PD 04-OCT-1994. 945283.
 PF 11-SEP-1992; US-945283.
 PR 11-SEP-1992; US-945283.
 PA (USDA) US SEC OF AGRIC.
 PI Cheung AK, Wesley RD;
 DR WPI: 94-316187/39.
 DR P-PSDB: R60620.
 PT New pseudorabies virus mutants for use in vaccine - having a
 PT deletion and/or insertion in the early protein 0 gene or large
 PT latency transcript gene
 PS Disclosure; Column 15-30; 43pp; English.
 CC Q73500 shows the Pseudorabies virus (PRV) large latency transcript
 CC (LNT). The basic sequence is derived from PRV strain InPh and PRV
 CC strain Ka. The LNT overlaps and is transcribed in the opposite
 CC orientation with respect to the EP0 (early polypeptide 0) and the
 CC immediately early gene (IE180). EP0 is nonessential for replicatio,
 CC LNT is the only gene expressed during PRV latency, and the IE180
 CC gene is absolutely necessary for PRV replication. However there are
 CC 2 copies of IE180 in the genome. It is expected that PRV lacking one
 CC of the IE180 copies is viable. Deletions in the non-overlapping
 CC regions of these 3 genes will generate single deletion routants,
 CC while deletions in overlapping regions will generate double deletion
 CC mutants. The invention is concerned with the construction of attenuated
 CC viruses which have a reduced ability to reactivate from latency. This
 CC can be achieved by functionally disabling the expression of the EP0
 CC gene, or by disrupting the synthesis of the LNT, or both.(See also
 CC Q73501 and R60620-24)
 CC Sequence 8438 BP; 1141 A; 2916 C; 3327 G; 1054 T;
 SQ

Query Match 1.3%; Score 41.6; DB 1; Length 8438;

Best Local Similarity 48.18; Pred. No. 1.5; Mismatches 159; Indels 1; Gaps 1;

QY 132 CCGGTCCAGGCTGGCGAGTCCGAGGCGGAGGAGGCGCGTGGTGGAGCAGATCC 191
DB 4305 CCAGCGCGGGCGCGCGGGCGCGCCCTCCCGGCTCCCGGCTCCCGGCTCC 4246
QY 192 AGAGCCGTGCGCCCCAGAACTGGCGGTCCCGCCCGCGTGCACCCCGCGGCATGCCAG 251
DB 4245 AGCCCGCGCAGCAGCAAGTCCGGTCCAGCAGCAAGTCCAGCTCCGGCAGTCCGGC 4186
QY 252 TTGCCCCCGCGCTGTGTAGGGCGCGCTCTCCATGATGAGGCTCATGGCGTCAACAA 311
DB 4185 CTCTCCGGCTCTCCGGGTAGCGGCTCCCGCCCGCGCGCGCGGACCCAGCGC-CCGA 4127
QY 312 GGAGGTGCCAAGGCGACATCGGCGGGGTGTCTCCCGCGCTGGAACTGGCCATCGA 371
DB 4126 GCGGGCAGAGAGAGCGCCGCGCGCGCGCGCGCGCGCGCGCGCGGAGGAGGA 4067
QY 372 GCAGATCGCAACAGTCACTCTGCGCCCTACTTCTCGACCTGCGGCTCTATGACAC 431
DB 4066 CGAGGGGTCTCCGGCTCGGCGCTCCGCGGGGACGCGCACGCGCCAGCGAGGA 4007
QY 432 GGAGTGGC 439
DB 4006 GGACGGG 3999

RESULT 15

T35233/C
ID T35233 standard; cDNA; 2823 BP.
AC T35233;
DT 05-DEC-1996 (first entry)
DE Natural killer lytic associated protein cDNA.
KW Natural killer lytic associated protein; NKLP;
KW cytotoxic T-lymphocyte; CTL; natural killer cell; antisense;
KW tumor; cancer; renal carcinoma; melanoma; herpes simplex virus;
KW hepatitis virus; infection; graft vs. host; autoimmune disease;
KW systemic lupus erythematosus; rheumatoid arthritis; gene therapy;
ds.
OS Homo sapiens.
FH Key
FT cds
FT signal_peptide 190..324
FT /tag= a
FT /tag= b
FT /note= "putative signal peptide"
FT mat_peptide 325..1950
FT /tag= c
FN W09626744-A1.
PD 06-SEP-1996.
PF 01-MAR-1996; U02736.
PR 02-MAR-1995; US-398008.
PA (UVAR-) UNIV ARKANSAS.
PI Kornbluth J.
DR WPI; 96-412588/41.
DR P-PSDB; R99256.
PT New human natural killer lytic associated protein - useful for
PT treating cancer, viral infections, graft vs. host or autoimmune
PT diseases.
PS Claim 2; Page 30-32; 65pp; English.
CC A cDNA clone (T35233) codes for a human natural killer lytic
CC associated protein (NKLP) (R99256). It was isolated by:
CC preparing an interferon-beta stimulated natural killer 3.3 cDNA
CC library; selecting cDNA from the library that was expressed at
CC elevated levels in interferon-beta stimulated 3.3 cells; and
CC sequencing the isolated clones. The cDNA can be used for prodn.
CC of commercial quantities of NKLP in transformed bacterial (esp.
CC E. coli), mammalian (esp. NK 3.3) or insect (esp. Sf9) host cells.
CC It is useful in gene therapy protocols to enhance the antitumor,
CC antiviral and antimicrobial activity of NK cells. Antisense
CC sequences (see also T35234-35) are useful therapeutically.
SQ Sequence 2823 BP; 653 A; 761 C; 807 G; 602 T;

Query Match 1.3%; Score 41.6; DB 1; Length 2823;
Best Local Similarity 44.7%; Pred. No. 0.96;
Matches 161; Conservative 0; Mismatches 199; Indels 0; Gaps 0;
QY 2778 CCAGCGTGGCTGCTCCCTCCAGTCCCTCCATCTCCACACAGCTTACCTCCCATCATCGG 2837
DB 559 CCAGGCACAGCGGACACTCCACTCTCTCCCGCGCGGCGCACCGCCCTCCGCGGCTCTCT 500
QY 2838 AGCGGTGGACGCGAGCTGTGTAGCCCTTGTAGCCCCCAGCCAGCCCCCGGCGCACAG 2897
DB 499 CATGCTGAACCCAGAGCTCCGCGCGCGCGCGCTCCGCGCTCGGCTCGGCGGCG 440
QY 2898 ACATGTGCCACCTCTCTCCGAGTGTGTCTCGGGCTGTAAAGGTGGAGGCTTGGGC 2957
DB 439 GCTCGCGGGGACGCGCTCGGGCGCGGCGCTTGGCGCGCGGCGCGGCGGCGG 380
QY 2958 CCGGGGCTTCCCGCTGACAGAACCACTGGGCGAGGGGTCTGTGCAGAAACACTGT 3017
DB 379 GCGGCGCGCAGCGCGGGGCGGCGCTCGGCTTGGCCCGGCGGCGGCGGCGGCG 320
QY 3018 CGGCTCTGCTGGGAGAGAGTGGGACCATGGCTGGCTCTCAGGACCACTCGGATGGC 3077
DB 319 GGGCGGAGGAGAGAGAGCTGTGCAAGGTGAGGCGCGGCGCGCGCGCTGCGGC 260
QY 3078 ACTCAGGTGGACAGGCGGCGGAGAGCTTGGCACCTGACCTCGAGCCTTATTGT 3137
DB 259 ACTTAGGTGGGTGGCGCGCATGTAGCGATGTGGAGCGGCGGCGGCGGAGTCTCTTCT 200

Search completed: March 16, 2000, 17:40:53
Job time: 3797 sec

